

APPROVED	Q.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

OBLON ET AL (703) 413-3000
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FIG. 1

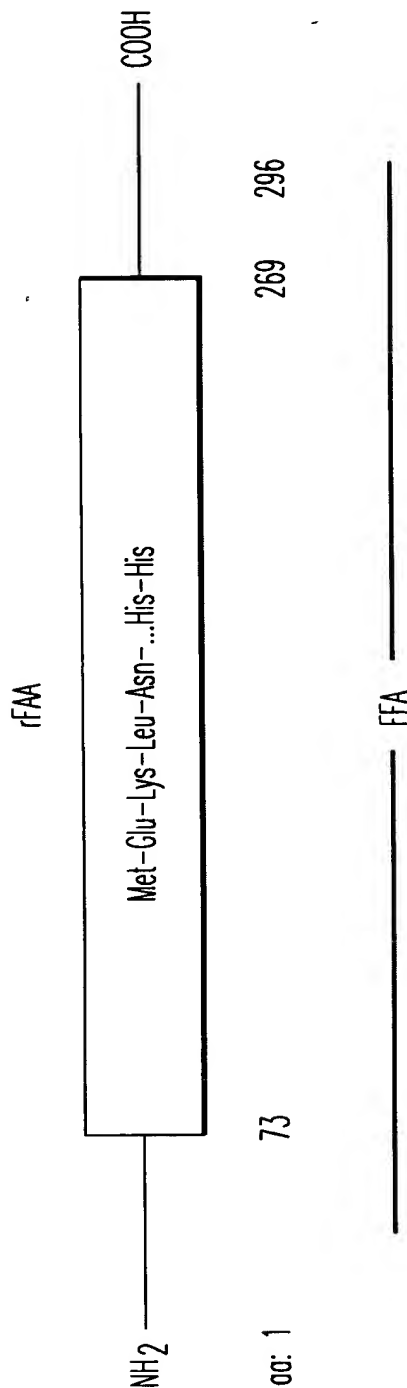


FIG. 1

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FIG. 2

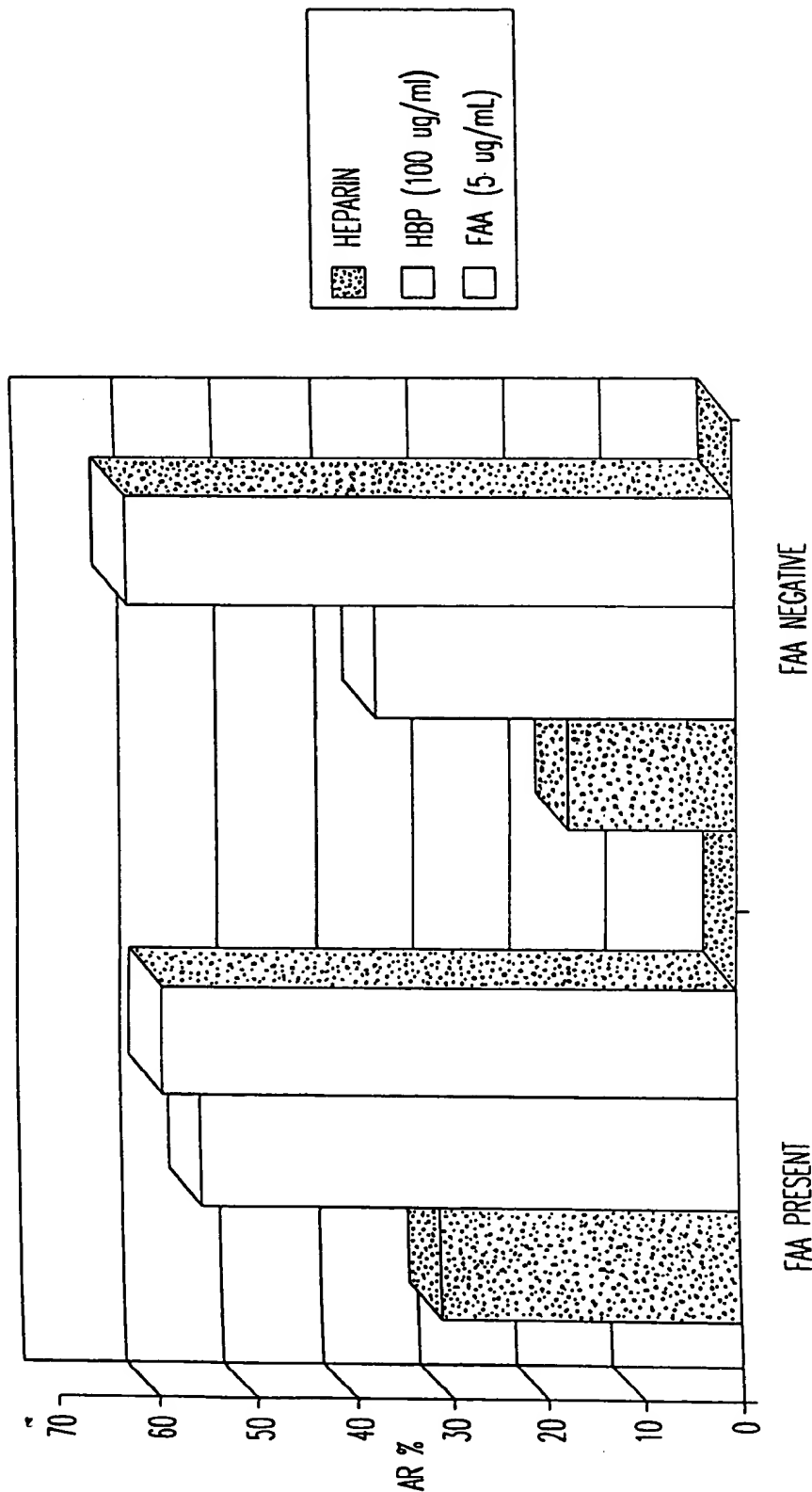
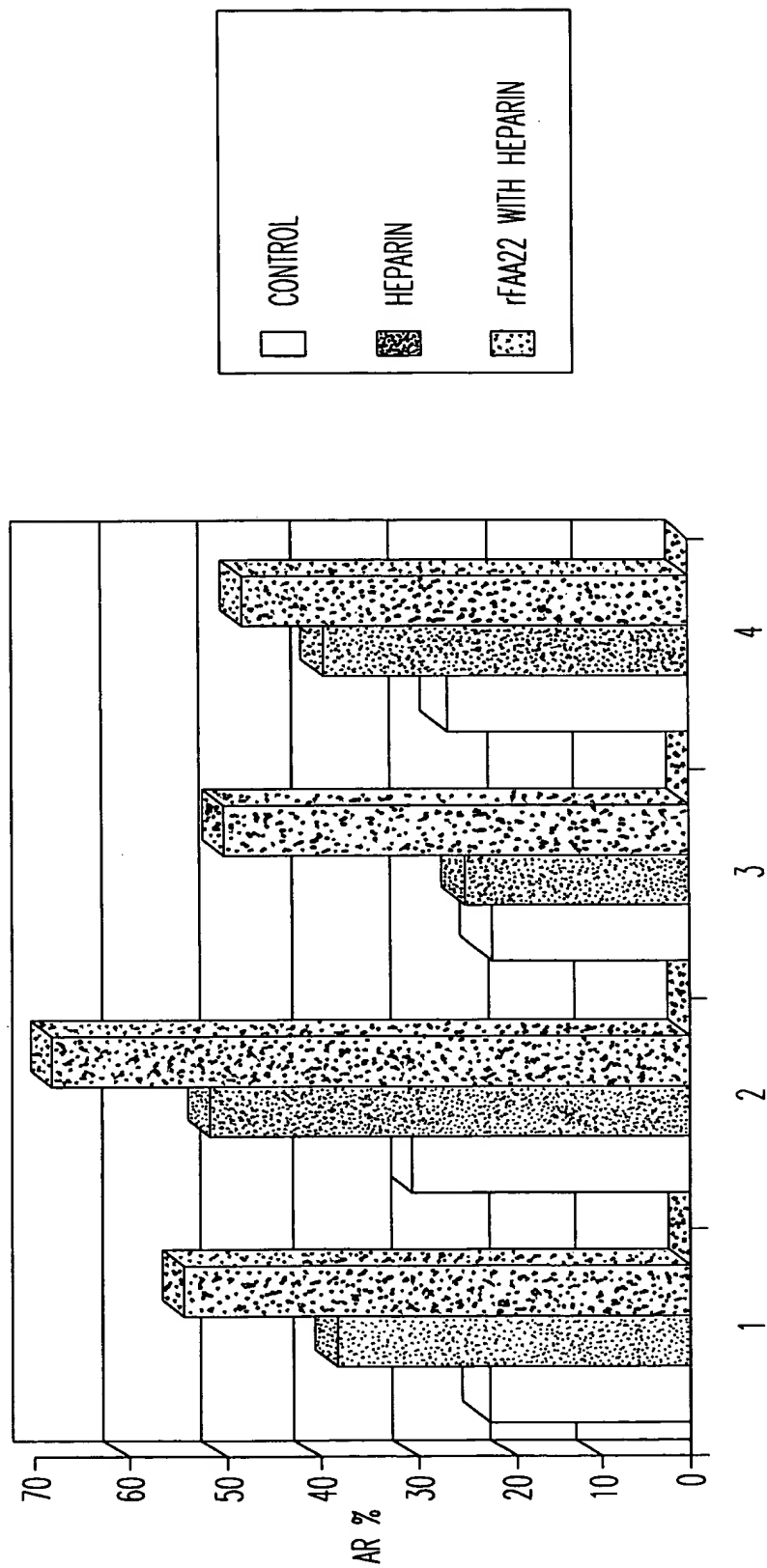


FIG. 2



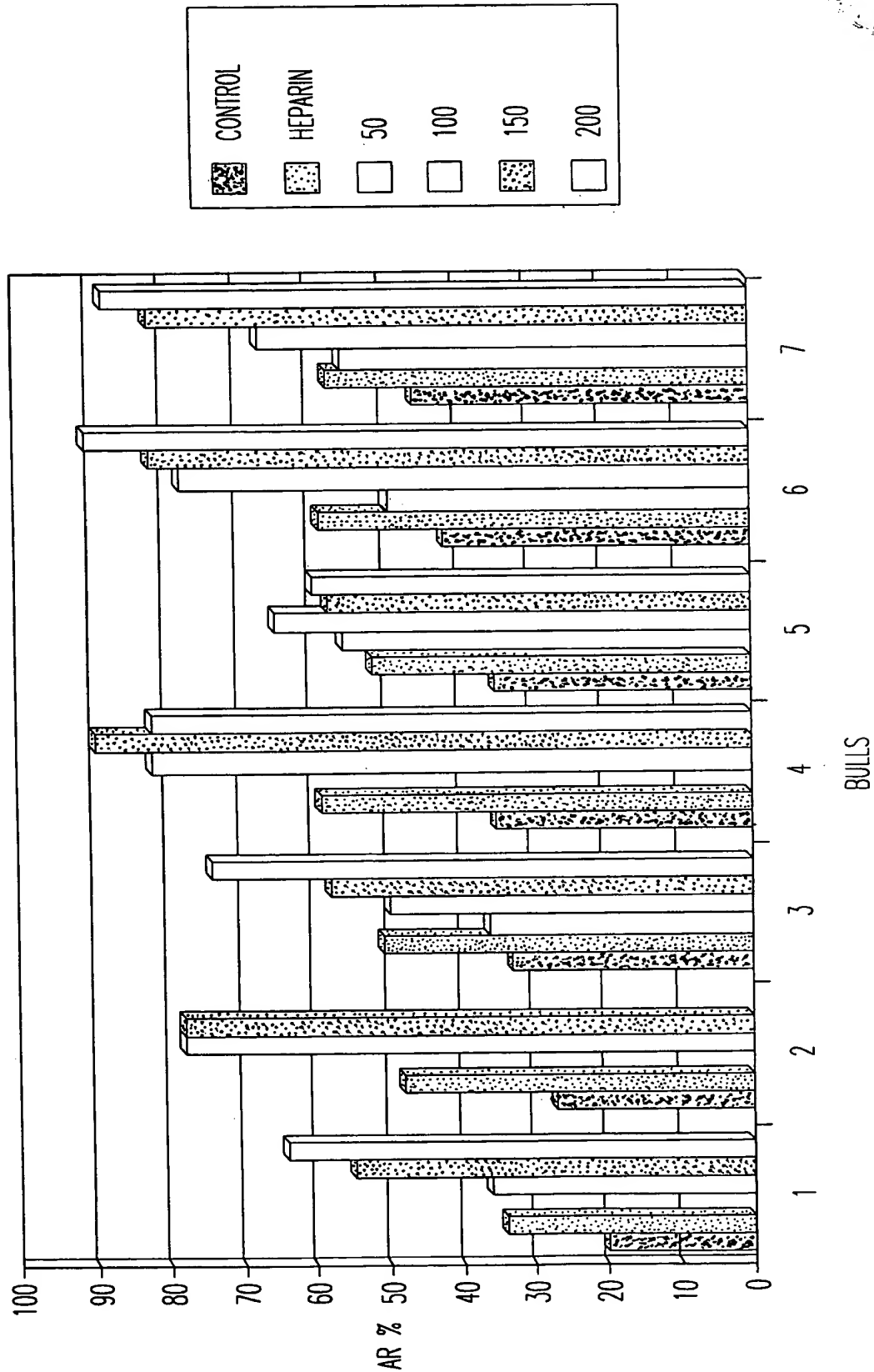
FIG. 3



BULL

FIG. 3

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1 ACAACAGGAT CTGCCCCATA CTGATGGAGA AGCTAACGG AAATTCAAGA
51 AAAGGCATAA CATACAATA TG TGATTAGC TCTCGCCTTG GAAGAAACAC
101 ATATAAGAA CAGTATGCCT TTCTCTATAA AGAAAAGCTA GTGTCTGTAA
151 AACAAAGCTA CCTCTACCAC GACTATCAGG CTGGAGACGC AGATGTGTTT
201 TCCAGGGAAC CCTTTGTGGT CTGGTTCCAG TCACCCTACA CCGCTGTCAA
251 GGACTTCGTG ATTGTCCCC TGACACCAC CCCTGAGACA TCCGTTAGAG
301 AGATTGATGA GCTGGCTGAT GTCTACACAG ATGTGAAACG TCGCTGGAAT
351 GCAGAGAATT TCATTTTCAT GGGTGACTTC AATGCTGGCT GCAGCTACGT
401 CCCCAGAAG GCCTGGAAGG ACATCCGCCT GAGGACGGAC CCAAGTTCG
451 TTTGGCTGAT CGGGGACCAA GAGGACACCA CGGTCAAGAA GAGCACAAAC
501 TCGCCTATG ACAGGATCGT GCTTAGAGGA CAAAATATTG TCAACTCTGG
551 TGGTCCTCAA TCAAACCTCG TCTTTGATTT CCAGAAAGCT TACAGGTTGT
601 CTGAATCGAA GGCCCTGGAT GTCAGCGACC ACTTTCCAGT TCATCATCAT
651 CATCATCATG AAGAACCATG A

FIG. 5

FIG. 5

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5' GAGAAGCTAAACGGAAATTCAAGAAAAGGCATAACATACTATGTGATTAGCTCTGGC 60
1 -----+-----+-----+-----+-----+-----+
a E K L N G N S R K G I T Y N Y V I S S R
CTTGAAGAAACACATATAAAGAACAGTATGCCTTTCTCTATAAAGAAAAGCTAGTGTCT
61 -----+-----+-----+-----+-----+-----+ 120
a L G R N T Y K E Q Y A F L Y K E K L V S -
GTAAACAAAGCTACCTCTACCACGACTATCAGGCTGGAGACGCAGATGTGTTTTCCAGG
121 -----+-----+-----+-----+-----+-----+ 180
a V K Q S Y L Y H D Y Q A G D A D V F S R -
GAACCCTTTGTGGTCTGGTTCCAGTCACCCCTACACCGCTGTCAAGGACTTCGTGATTGTC
181 -----+-----+-----+-----+-----+-----+ 240
a E P F V V W F Q S P Y T A V K D F V I V -
CCCCTGCACACCACCCCTGAGACATCCGTTAGAGAGATTGATGAGCTGGCTGATGTCTAC
241 -----+-----+-----+-----+-----+-----+ 300
a P L H T T P E T S V R E I D E L A D V Y -
ACAGATGTGAAACGTCGCTGGAATGCAGAGAATTTTCATTTTCATGGGTGACTTCAATGCT
301 -----+-----+-----+-----+-----+-----+ 360
a T D V K R R W N A E N F I F M G D F N A -
GGCTGCAGCTACGTCCCCAAGAAGGCCTGGAAGGACATCCGCCTGAGGACGGACCCCAAG
361 -----+-----+-----+-----+-----+-----+ 420
a G C S Y V P K K A W K D I R L R T D P K -
TTCGTTTGGCTGATCGGGGACCAAGAGGACACCACGGTCAAGAAGAGCACAACTGCGCC
421 -----+-----+-----+-----+-----+-----+ 480
a F V W L I G D Q E D T T V K K S T N C A -
TATGACAGGATCGTGCTTAGAGGACAAAATATTGTCAACTCTGGTGGTCCTCAATCAAAC
481 -----+-----+-----+-----+-----+-----+ 540
a Y D R I V L R G Q N I V N S G G P Q S N -
CTCGTCTTTGATTTCCAGAAAGCTTACAGGTTGTCTGAATCGAAGGCCCTGG 3'
541 -----+-----+-----+-----+-----+-----+ 592
a L V F D F Q K A Y R L S E S K A L -

FIG. 6